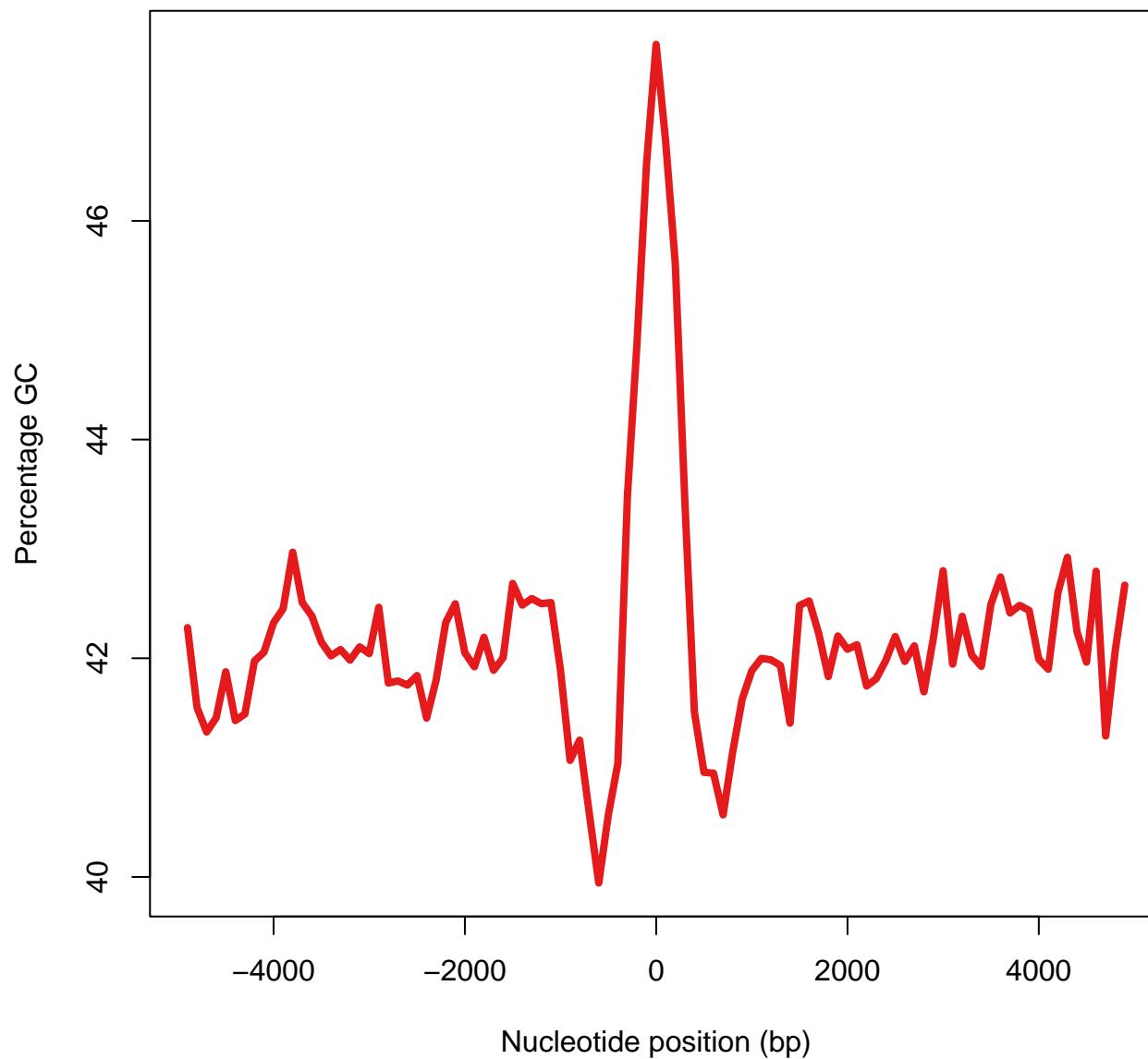
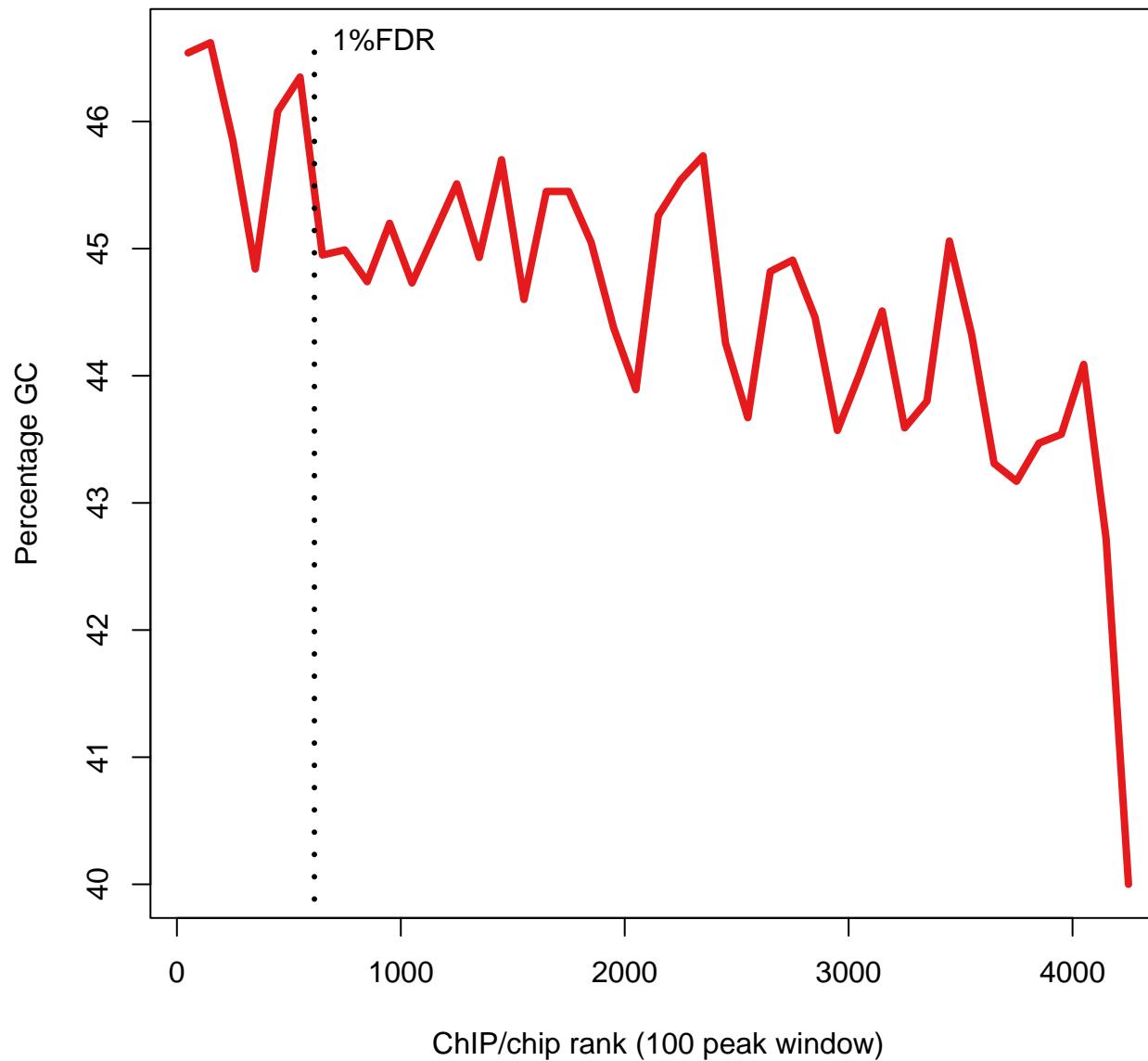


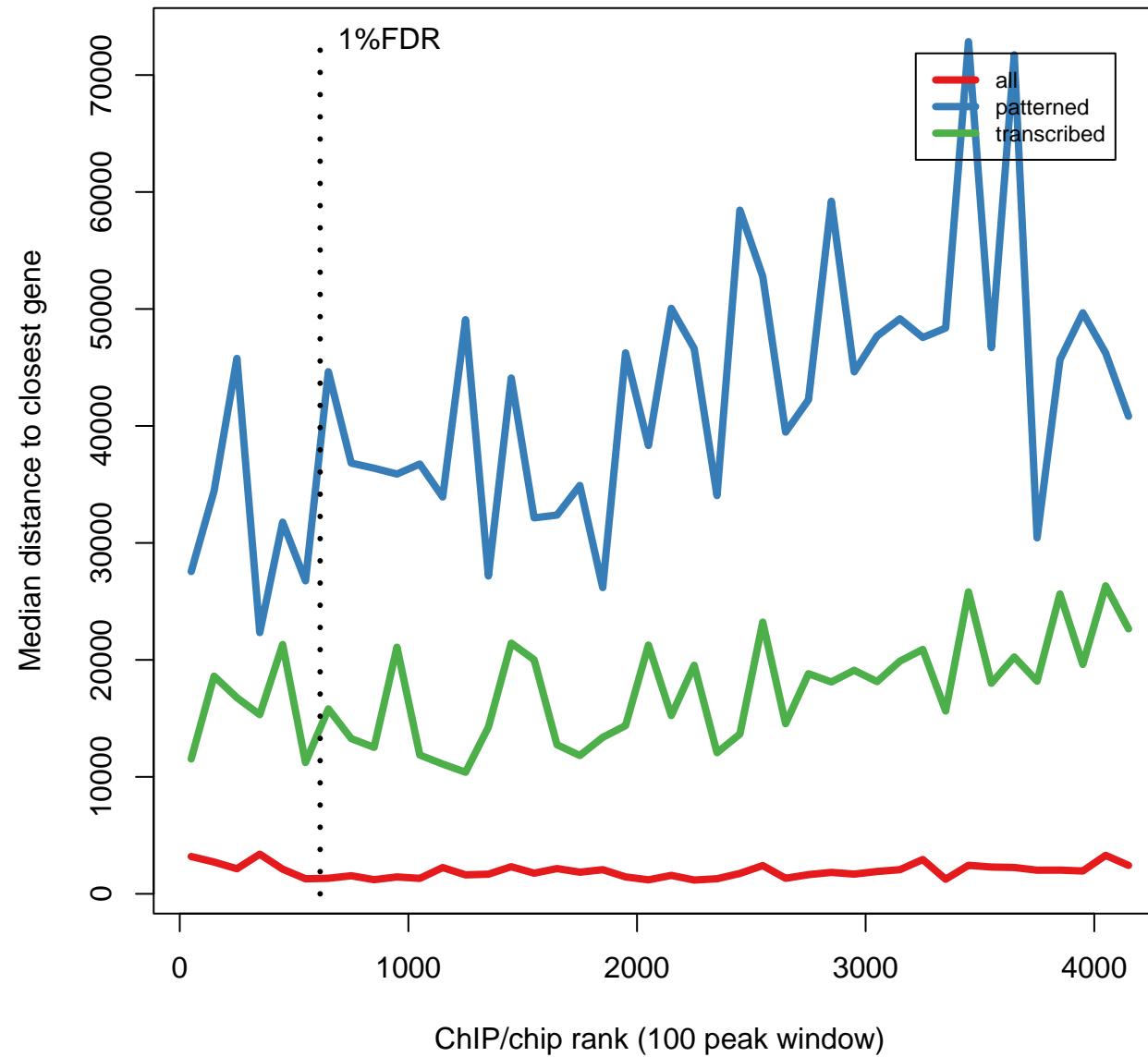
## **hkb\_2 base composition across peaks**



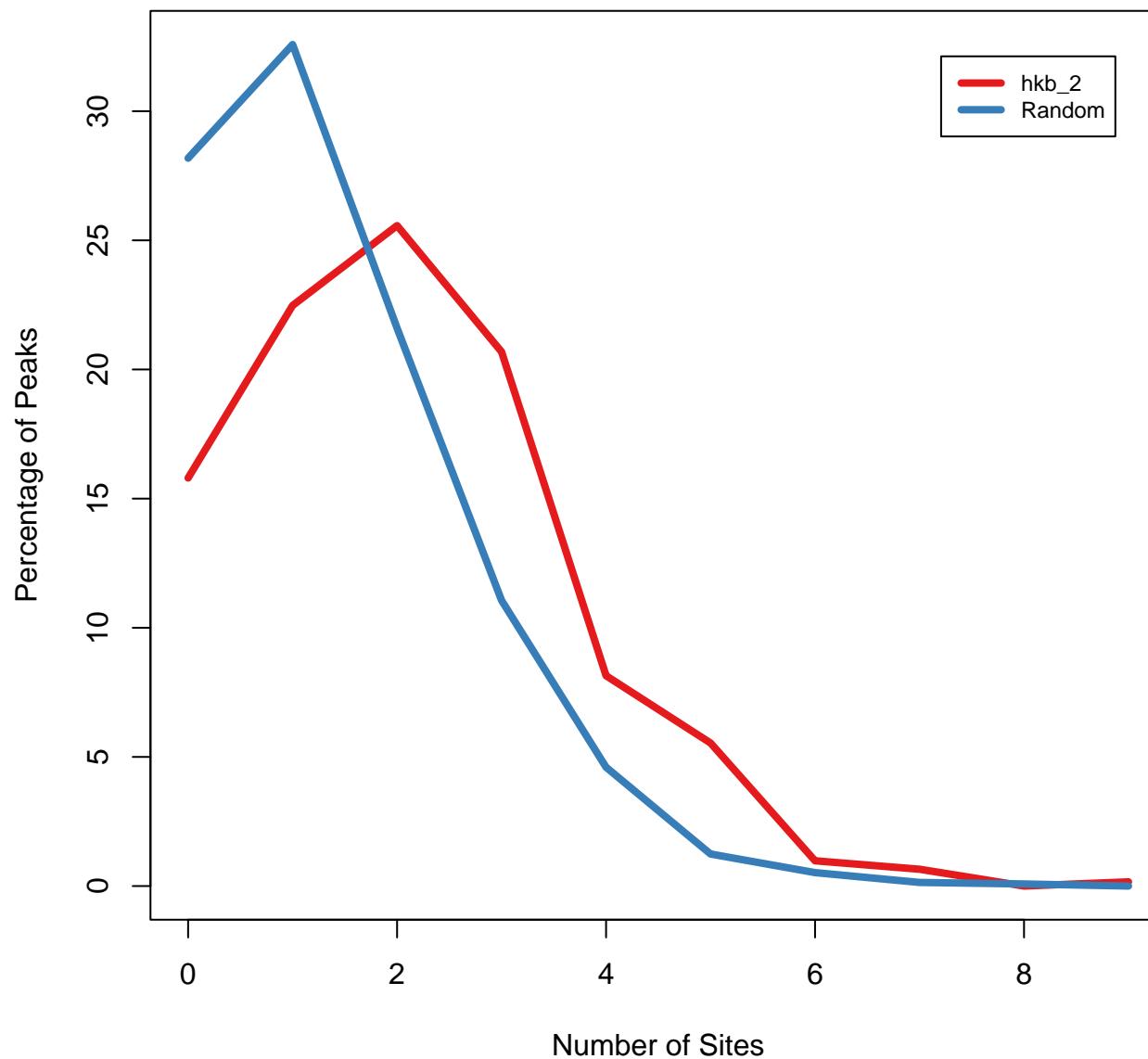
### **hkb\_2 base composition down ranks**



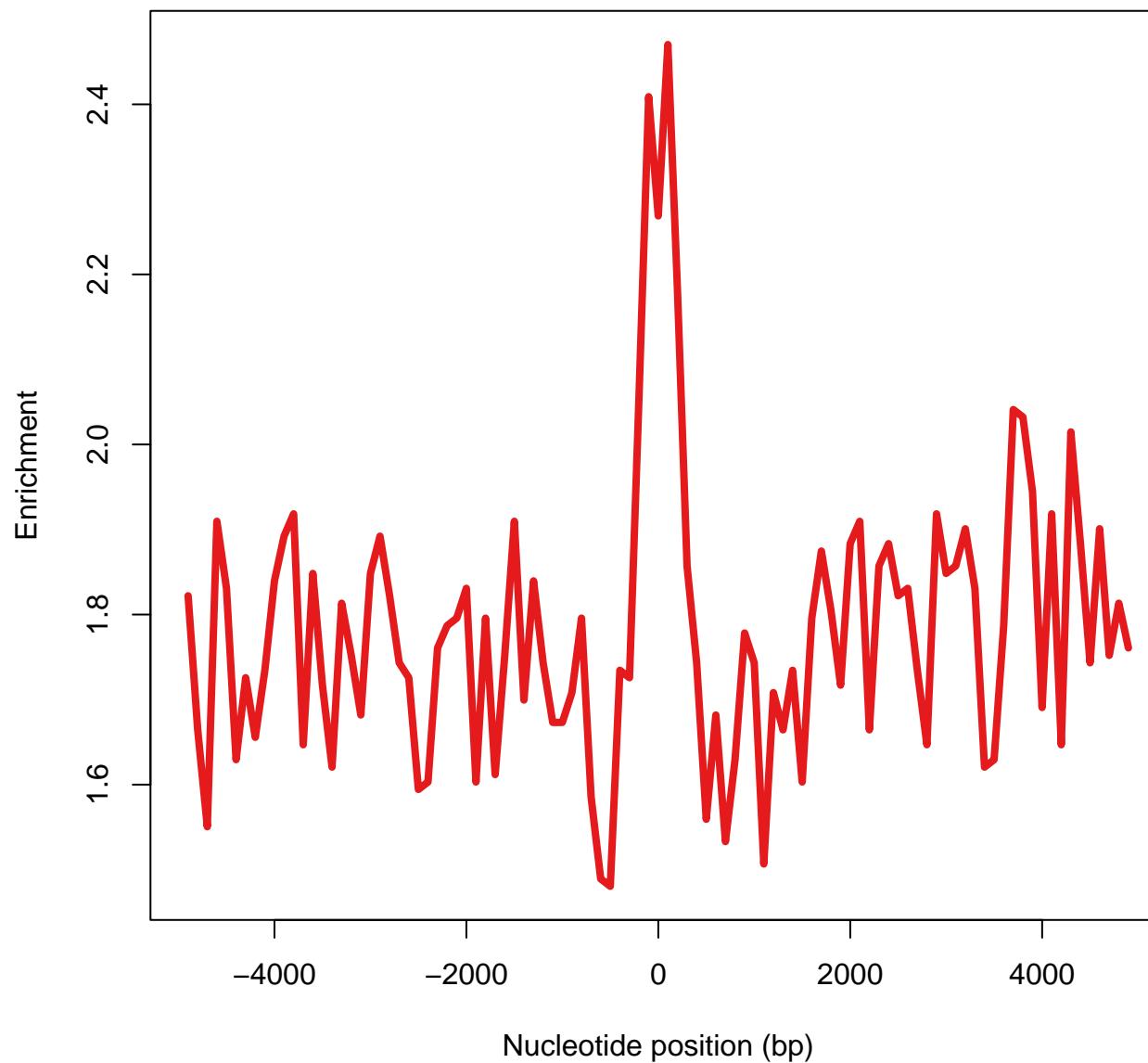
## hkb\_2 median distance to genes



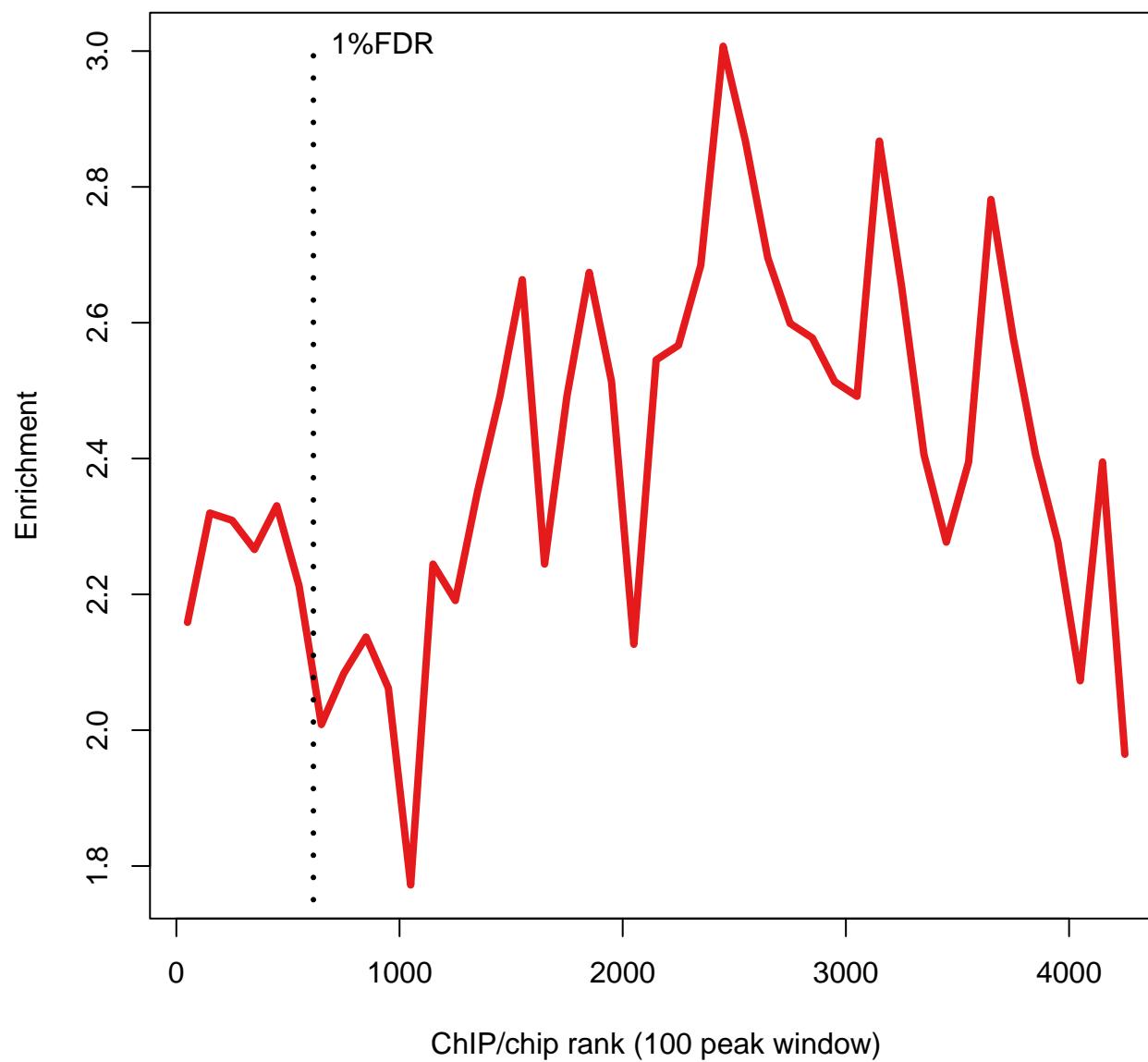
### **hkb\_2 distribution of number of sites**



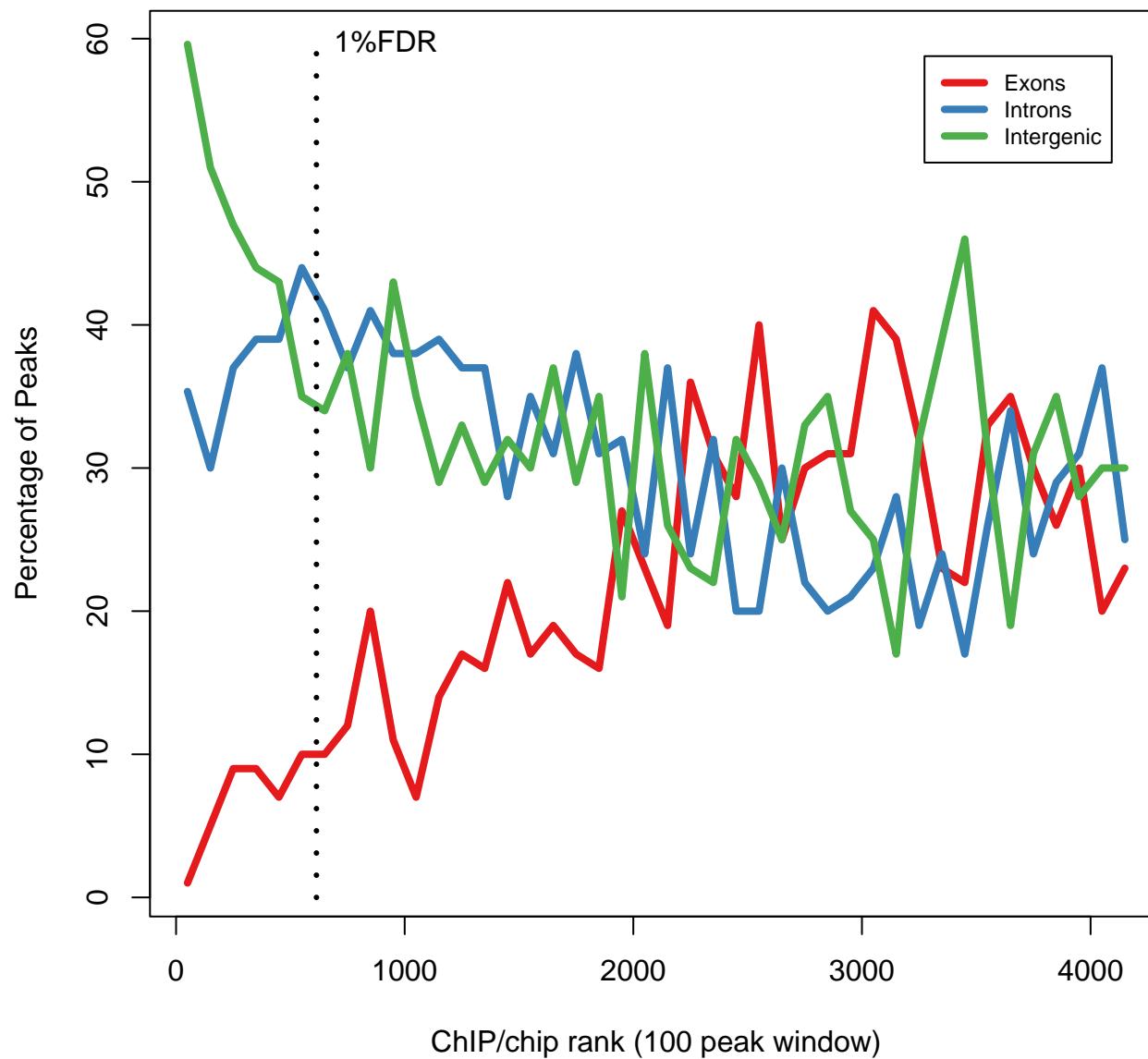
### **hkb\_2 PWM enrichment across peaks**



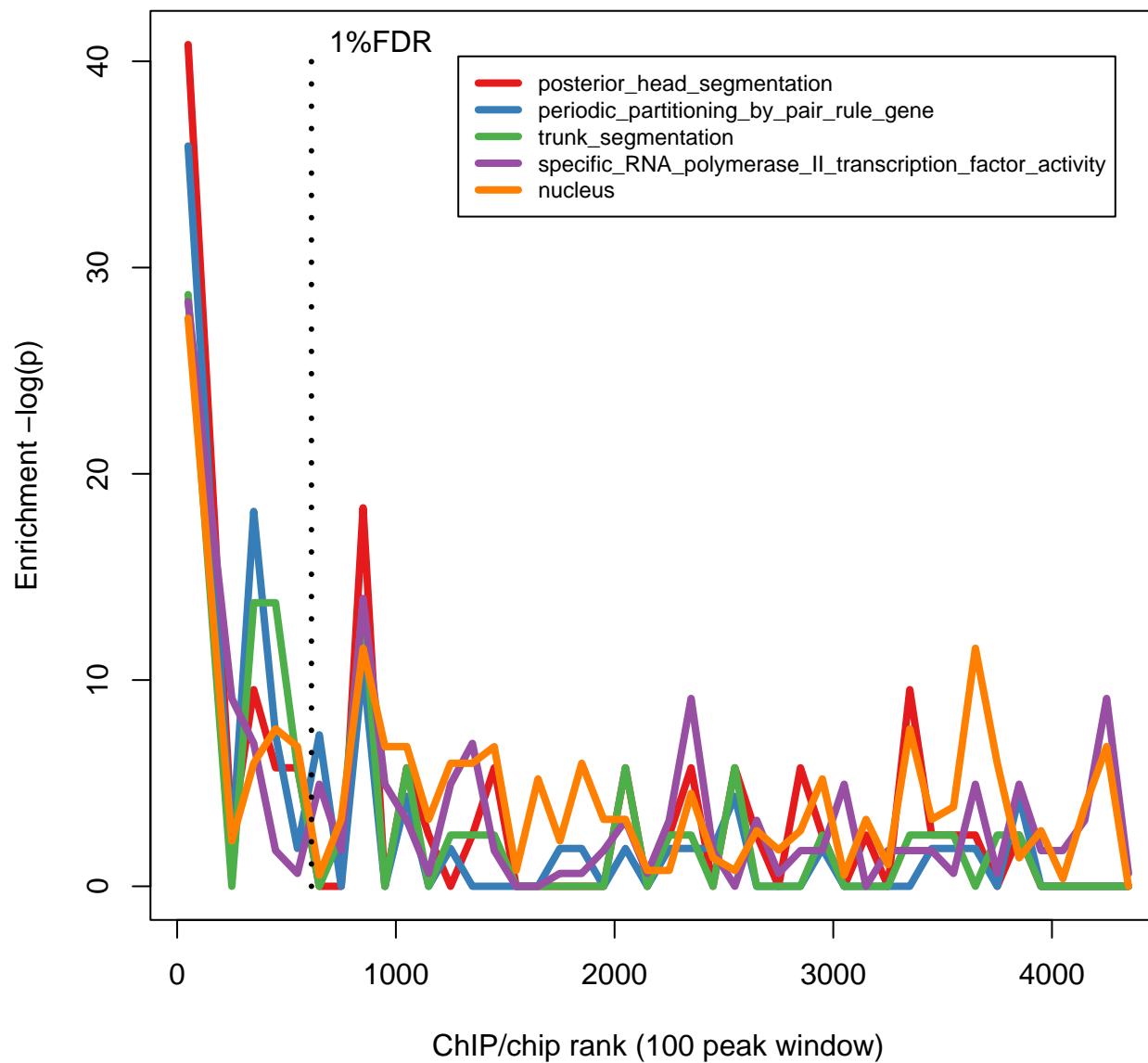
### **hkb\_2 PWM enrichment down ranks**



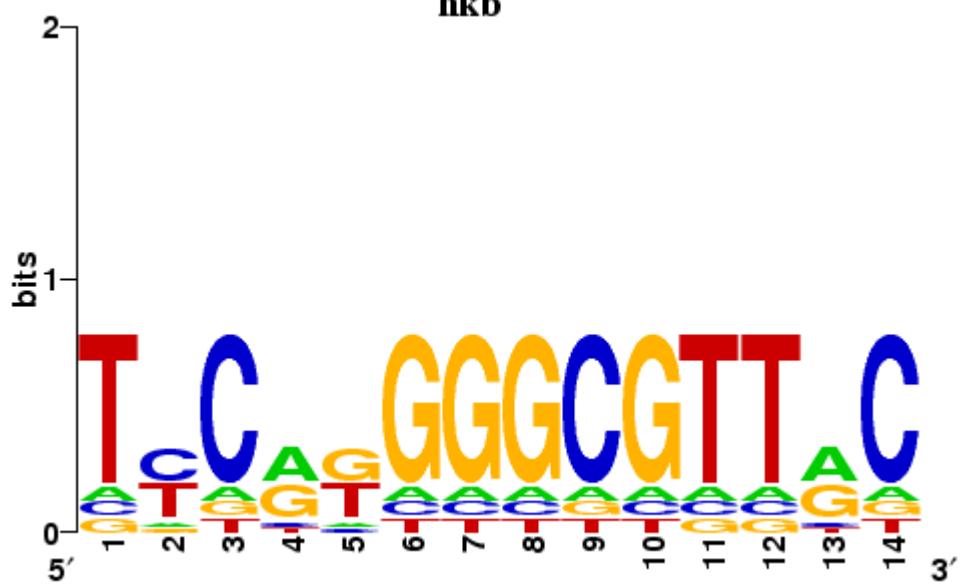
## **hkb\_2 genomic location**



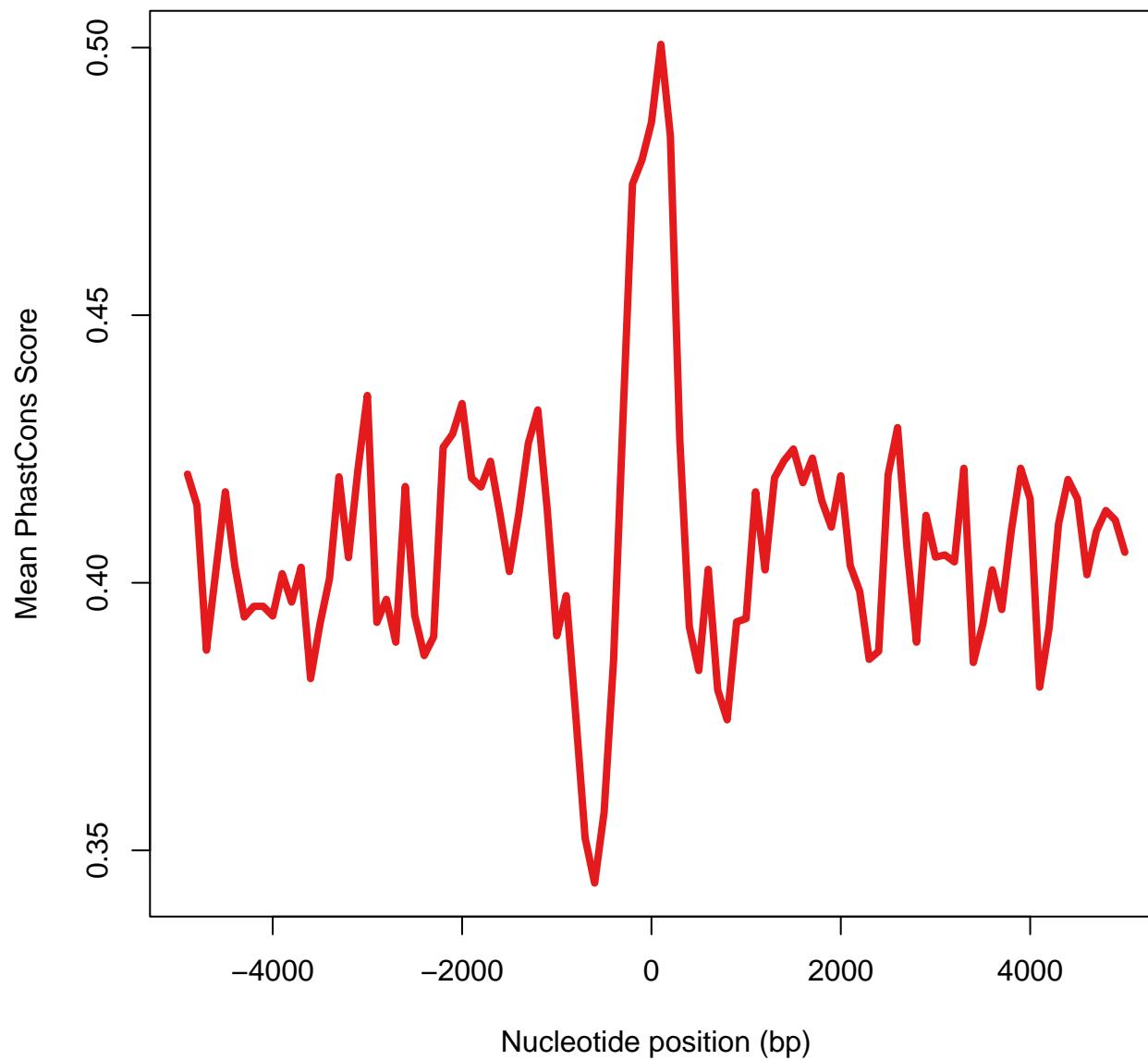
## hkb\_2 GO term enrichment ( $-\log P$ )



**hkb**



### **hkb\_2 mean PhastCons across peaks**



## **hkb\_2 mean PhastCons down ranks**

